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(57) Abstract

The labeling of biologically important molecules via a bifunctional chelator can result in the formation of isomers or multiple species, which can have significant impact on the biological properties of the radiopharmaceutical. For receptor–based radiopharmaceuticals, the target uptake is largely dependent on the receptor binding affinity of the targeting molecule and the blood clearance of the labeled molecule, which is determined by the physical properties of both the targeting molecule and the metal chelate. Hence, the presence of isomers for the metal chelate can have significant impact on the radiopharmaceutical. Therefore, in the development of a radiopharmaceutical or metallodrug, it is necessary to separate the isomers and evaluate the biological activities of each individual isomer. It would therefore be desirable to develop chelators that predominately form only a single stereoisomeric species upon coordination to a metal center. Disclosed herein are chelators that form a mixture enriched for a single stereoisomeric species upon coordination to a metal center.

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PEPTIDE CHELATORS THAT PREDOMINATELY FORM A SINGLE STEREOISOMERIC SPECIES UPON COORDINATION TO A METAL CENTER

Technical Field

This invention relates to chelators that form a mixture enriched for a single stereoisomeric species upon coordination to a metal center.

Background of the Invention

The current interest in radiolabeling biologically important molecules (proteins, antibodies, and peptides) with ^{99m}Tc stems from the desire to develop a target specific diagnostic radiopharmaceutical.¹⁻¹⁰ The advantages of using ^{99m}Tc in diagnostic nuclear medicine are well known¹¹⁻¹⁵ and a number of techniques have been developed for the ^{99m}Tc labeling of biologically important molecules.¹⁶⁻²⁰

One obvious approach is to coordinate a ^{99m}Tc metal directly with the targeting molecule. This approach is known as the direct labeling method and it involves the use of a reducing agent to convert disulfide linkages into free thiolates, which then bind to the ^{99m}Tc metal. A major disadvantage of this method is the lack of control over the coordination of the ^{99m}Tc metal and the stability of the resulting metal complex. In addition, the lack of suitable or accessible coordination sites in some proteins and peptides exclude direct labeling as a viable technique.

Two common alternatives to direct labeling are the final step labeling method and the pre-formed chelate method. Both techniques involve the use of a bifunctional chelator, which provides the site of ^{99m}Tc coordination. The difference between the two methods lies in the order in which the ^{99m}Tc complex is formed. In the final step labeling method, complexation occurs after the chelator has been attached onto the targeting molecule. With the pre-formed chelate method, the ^{99m}Tc complex is initially prepared and purified before being attached to the targeting molecule. In both methods, the bifunctional chelator must coordinate to ^{99m}Tc to form a complex that is stable *in vivo* and the chelator must have an active moiety that can react with a functional group on the targeting molecule.

A number of bifunctional chelators have been used in the labeling of proteins, peptides and monoclonal antibodies.^{2, 9, 10, 17, 21-28} Depending on the chelator, the

labeling of biologically important molecules with bifunctional chelators often results in the formation of multiple species or isomeric complexes. An example is the 99mTc labeling of molecules using the hydrazinonicotinamide (HYNIC) system. Since the HYNIC group can only occupy one or two sites of Tc coordination, co-ligands are Glucoheptonate²⁹⁻³⁰, required to complete the coordination sites. tris(hydroxymethyl)methylglycine (tricine)²⁵, ethylenediamine-N, N'-diacetic acid (EDDA)⁹, water soluble phosphines²⁵ [trisodium triphenylphosphine-3,3',3"trisulfonate (TPPTS), disodium triphenylphosphine-3,3'disulfonate (TPPDS), and sodium triphenylphosphine-3-monosulfonate (TPPMS)] and polyamino polycarboxylates9 have all been used as co-ligand in the HYNIC system.

It has been clearly shown that the Tc-99m labeling of molecules via the HYNIC/coligand system produces multiple species, which is due to the different coordination modalities of the hydrazine moiety and the co-ligands. The number of species, the type, the stability and the properties of the species vary greatly from one co-ligand to another. In the labeling of chemotactic peptides using the HYNIC system, the nature of the co-ligand also greatly affects the biodistribution of the labeled peptide.³¹

Another example of a bifunctional chelator producing multiple species is dithiosemicarbazone (DTS) system. It has been shown that the DTS bifunctional chelator produces at least four complexes with technetium.³² Two of the complexes are known to be charged; hence they have different biodistribution from the uncharged species.

As in the development of a pharmaceutical based on organic molecules, the stereochemistry or isomerism of a metal complex is very important in the development of a radiopharmaceutical or metallodrug. It is well known that isomers can have different lipophilicities, biodistribution patterns, and biological activities. An example of this is the ^{99m}Tc complex of 3,6,6,9-tetramethyl-4,8-diazaundecane-2,10-dione dioxime (^{99m}Tc-d,l-HMPAO or Ceretec), which is a cerebral perfusion imaging agent. ^{14,33-35} Though ^{99m}Tc-d,l-HMPAO is active, it has been shown that the meso analogs of the ^{99m}Tc HM-PAO^{14,36} complex and the ^{99m}Tc complex of 3,3,9,9-tetramethyl-4,8-diazaundecane-2,10-dione dioxime^{14,37} (PnAO) do not possess the properties necessary for use as a cerebral perfusion imaging agent.

A type of Tc and Re coordination modality common in Tc and Re radiopharmaceuticals is the coordination of a tetradentate N_{4-x}S_x chelator to a metal oxo moiety to form a square pyramidal or octahedral metal oxo complex. A host of bifunctional chelators have been developed based on the tetradentate N_{4-x}S_x coordination motif. Examples include N₄ propylene amine oxime³⁸, N₃S triamide thiols $^{9, 39-43}$, N_2S_2 diamide dithiols $^{9, 44-46}$, N_2S_2 monoamide monoaminedithiols $^{47-49}$ and N₂S₂ diamine dithiols⁵⁰⁻⁵⁵. Functionalization of the chelator backbone enables these chelators to be attached to biologically interesting molecules. The labeling of these bifunctional chelators with TcO3+ or ReO3+ often produces isomers or epimers. 39-43, 46-55 The isomers or epimers (syn and anti) arise from the configuration of the metal oxo group relative to the functional group on the chelator backbone. It has been clearly shown that the biodistribution and biological activity of the syn and anti isomers are often different. 39-43, 46, 56 The Tc complex of mercaptoacetylglycylglycine (MAG₃), a renal imaging agent, exists in the syn and anti isomers. The biological activities of the syn and anti isomers are known to be different.^{39,40} The syn and anti isomers of the Tc complex of 2,3-bis(mercaptoacetamide)propanoate (map) wwere also shown to have different biological activities.46 It was reported that, in humans, 58% of the syn isomer was excreted at 30 minutes as compared to only 19 % of the anti isomer. Another example of the isomers exhibiting a difference in biological behaviour is the 99mTc labeled diamino dithiol piperidine conjugates, which were investigated as brain perfusion imaging agents. It was shown that the two isomeric complexes exhibit widely disparate brain uptake. 55 At 2 minute post-administration in rats, uptake of the anti isomer in the brain was 1.08 % dose/g, while the uptake of the syn isomer was 2.34 % dose/g. The brain/blood ratio at 2 minute post-administration was 2.09 for the anti isomer and 5.91 for the syn isomer.

The peptide dimethylglycine-L-serine-L-cysteine-glycine is a bifunctional chelator that can be used to label biologically important molecules. It has been shown that dimethylglycine-L-serine-L-cysteine-glycine coordinates to TcO³⁺ and ReO³⁺ via a monoamine diamide monothiol coordination modality. The resulting Tc and Re complexes exist as two isomers; the serine CH₂OH side chain is in the *syn* and *anti* conformations with respect to the metal oxo bond. The presence of the *syn* and *anti* isomers is very evident from the NMR spectral data. In the ¹H NMR spectrum of the

Re complex, there were two pairs of singlets associated with the nonequivalent methyl groups in the dimethylglycine residue. Each pair of singlets corresponded to either the *syn* or *anti* isomers. The presence of the two isomers is clearly evident from the NMR data. In the coordination of dimethylglycine-L-isoleucine-L-cysteine-glycine (RP349) to ReO^{3+} , two isomers (*syn* and *anti*) were also observed. The 99m Tc labeling of RP294 and RP349 produced *syn* and *anti* isomers; two peaks were observed in the HPLC using the radiometric detector. The 99m Tc labeling of biotin with dimethylglycine-L-lysine-L-cysteine-NH₂ (RP332) also produced *syn* and *anti* isomers; two peaks were observed in the HPLC. These results are consistent with the coordination of other tetradentate $N_{4-x}S_x$ chelators to TcO^{3+} and ReO^{3+} . $^{9,39-55}$

The labeling of biologically important molecules via a bifunctional chelator can result in the formation of isomers or multiple species, which can have significant impact on the biological properties of the radiopharmaceutical. For receptor-based radiopharmaceuticals, the target uptake is largely dependent on the receptor binding affinity of the targeting molecule and the blood clearance of the labeled molecule, which is determined by the physical properties of both the targeting molecule and the metal chelate. Hence, the presence of isomers for the metal chelate can have significant impact on the radiopharmaceutical. Therefore, in the development of a radiopharmaceutical or metallodrug, it is necessary to separate the isomers and evaluate the biological activities of each individual isomer. It would therefore be desirable to develop chelators that predominately form a single stereoisomeric species upon coordination to a metal center.

Summary of the Invention

Chelators and chelator-targeting molecule conjugates are provided that form a mixture with a predominant stereoisomeric species upon coordination to a metal center.

According to an aspect of the invention, there is provided a chirally pure compound of the formula I:

$$R^{1}$$
 R^{2}
 R^{3}
 R^{4}
 R^{5}
 R^{6}
 R^{7}
 R^{8}
 R^{9}
 R^{10}
 R^{10}
 R^{10}
 R^{10}
 R^{10}

wherein

R¹ is a linear or branched, saturated or unsaturated C₁₋₄alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents selected from halogen, hydroxyl, amino, carboxyl, C₁₋₄alkyl, aryl and C(O)R¹⁰;

R² is H or a substituent defined by R¹;

- R¹ and R² may together form a 5- to 8-membered saturated or unsaturated heterocyclic ring optionally substituted by one or more substituents selected from halogen, hydroxyl, amino, carboxyl, oxo, C₁₋₄alkyl, aryl and C(O)Z;
- R³, R⁴ and R⁵ are selected independently from H; carboxyl; C_{1.4}alkyl; C_{1.4}alkyl substituted with a substituent selected from hydroxyl, amino, sulfhydryl, halogen, carboxyl, C_{1.4}alkoxycarbonyl and aminocarbonyl; an alpha carbon side chain of a D- or L-amino acid other than proline; and C(O)R¹⁰;

 R^6 is selected from a group consisting of :

- i) an optionally substituted 3- to 6-membered heterocylic or carbocylic ring,
- ii) a compound of the following formula:

wherein R^{11} , R^{12} and R^{13} are independently selected from H, linear or branched, saturated or unsaturated C_{1-6} alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R^{11} , R^{12} and R^{13} is not H;

iii) a compound of the following formula:

wherein R¹⁴ and R¹⁵ are independently selected from H, linear or branched, saturated or unsaturated C_{1.6}alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents (; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R¹⁴ and R¹⁵ is not H;

and iv) a compound of the following formula:

wherein X is selected from O or S and R¹⁶ is selected from linear or branched, saturated or unsaturated C₁₋₆alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, and an optionally substituted 3- to 6-membered heterocylic or carbocylic ring;

R⁷ and R⁸ are selected independently from H; carboxyl; amino; C₁₋₄alkyl; C₁₋₄alkyl substituted by a substituent selected from hydroxyl, carboxyl and amino; and C(O)R¹⁰;

R9 is selected from H and a sulfur protecting group; and

R¹⁰ is selected from hydroxyl, alkoxy, an amino acid residue, a linking group and a targeting molecule.

According to another aspect of the invention, there is provided a chirally pure compound of the formula II:

wherein

Ra is selected from H and a sulfur protecting group;

R^b, R^c R^d, R^f and R^g are selected independently from H; carboxyl; C₁₋₄alkyl; C₁₋₄alkyl substituted with a substituent selected from hydroxyl, amino, sulfhydryl, halogen, carboxyl, C₁₋₄alkoxycarbonyl and aminocarbonyl; an alpha carbon side chain of a D- or L-amino acid other than proline; and C(O)R^h;

Re is selected from a group consisting of:

an optionally subsituted 3- to 6-membered heterocylic or carbocylic ring;

and

wherein R^i , R^j and R^k are independently selected from H, linear or branched, saturated or unsaturated C_{1-6} alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R^i , R^j and R^k is not H;

and

wherein R^1 and R^m are independently selected from H, linear or branched, saturated or unsaturated C_{1-6} alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R^1 and R^m is not H;

and

wherein X is selected from O or S and Rⁿ is selected from linear or branched, saturated or unsaturated C_{1.6}alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, and an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; and

R^h is selected from hydroxyl, alkoxy, an amino acid residue, a linking group and a targeting molecule.

According to another aspect of the invention, the chelator-targeting molecule conjugates are provided in combination with a diagnostically useful metal or an oxide or nitride thereof.

According to another aspect of the present invention, there is provided a method of imaging a site of diagnostic interest, comprising the step of administering a diagnostically effective amount of a composition comprising a chelator-targeting molecule conjugate which is complexed to a diagnostically useful metal or an oxide or nitride thereof.

Detailed Description of the Invention

In the coordination of dimethylglycine-t-butylglycine-cysteine-glycine to TcO^{3+} and ReO^{3+} , a single isomer was observed. A single pair of singlets associated with the methyl groups in the dimethylglycine residue was observed. The ^{99m}Tc labeling of dimethylglycine-L-t-butylglycine-L-cysteine-glycine (RP455) and of dimethylglycine-D-t-butylglycine-L-cysteine-glycine (RP505) produced a single peak as observed in the HPLC using the radiometric detector. This was an unexpected result and contrasted with what was observed in the Tc and Re oxo complexes of other tetradentate $N_{4-x}S_x$ chelators, $^{9,39-55}$ which existed as the syn and anti isomers.

The presence of a sterically bulky group in the side chain of the peptidic chelator caused the formation of a single isomeric metal complex. In the cases of dimethylglycine-L-lysine-L-cysteine and dimethylglycine-L-serine-L-cysteine-glycine, there was insufficient bulk to cause one isomer to be preferred over another; hence the ratio of the *syn* and *anti* isomers was approximately 1:1.

In the case of dimethylglycine-L-isoleucine-L-cysteine, a more sterically bulky CH(CH₃)-CH₂-CH₃ group was introduced into the peptidic backbone. This additional bulk caused the ratio of the *syn* and *anti* isomers to be 3:1; hence, one isomer was more favored over the other. In the case of dimethylglycine-*t*-butylglycine-cysteine-glycine, the incorporation of the C(CH₃)₃ group introduced sufficient bulk into the peptide to cause one of the isomers to be completely favored over the other; hence, a single isomeric metal complex was observed.

Molecular modeling with Quanta Charm indicated that the *syn* isomer was favoured because in the *anti* isomer there was steric interaction between the bulky side group and the oxygen atoms of the adjacent amide groups. For example, the dihedral angles of the beta carbon of serine with the backbone of the chelate in the *anti* isomer of the Re complex of dimethylglycine-L-serine-L-cysteine-glycine (ReORP414) were –

27.39° (O-C-C-C) and 8.35° (C-C-N-C). The corresponding dihedral angles for the *anti* isomer of the Re complex of dimethylglycine-L-t-butylglycine-L-cysteine-glycine (ReORP455) were -11.95° and -6.87°. The difference of about 15° for each angle was a result of the shift of the amide oxygen atoms and the side group atoms of ReORP455 to a position of least contact. The shift of atomic positions induced some strain on the chelate system and therefore lessened its stability.

Molecular modeling of each of the Re complexes of the peptides was in agreement with experimental results. Molecular modeling of the Re complex of dimethyglycine-L-serine-L-cysteine-glycine showed the two isomers possessing thermodynamic potential energies of -67.02 and -68.37 kcal/mole. There was only a small difference in the energy of the two isomers. There was no preferred isomer for the Re complex and both the *syn* and *anti* isomers were observed at an approximate ratio of 1:1. Molecular modeling of the Re complex of dimethylglycine-L-lysine-L-cysteine showed a difference between the thermodynamic potential energies of the two isomers to be approximately 1 kcal/mole. There was again only a small difference in the energy of the two isomers; hence, both the *syn* and *anti* isomers would be observed.

In the case of dimethylglycine-L-isoleucine-L-cysteine-glycine, a more bulky side chain was incorporated into the peptidic backbone. Molecular modeling of the Re complex of the dimethylglycine-L-isoleucine-L-cysteine-glycine showed one of the isomers having a potential energy that was approximately 3 kcal/mole lower than the energy of the other isomer. There was now a greater difference in the energies and there was a slight preference for one isomer over the other. Accordingly, the observed experimental ratio of the two isomers was 3:1.

In the case of dimethylglycine-L-t-butylglycine-L-cysteine-glycine, molecular modeling of the Re complex showed the difference in the potential energies of the two isomers to be approximately 6.5 kcal/mole. With the Re complex of dimethylglycine-D-t-butylglycine-L-cysteine-glycine, the difference in the energies of the two isomers was about 8.5 kcal/mole. One isomer was significantly preferred over the other; hence, only a single isomer was observed for the Re and Tc complexes.

Molecular modeling of the Re complex of mercaptoacetyl-L-t-butylglycine-glycine-glycine showed that the *syn* and *anti* isomers of the complex with a energy difference of 7.4. The metal complexes of mercaptoacetyl-L-t-butylglycine-glycine-glycine preferred one isomer over the other and would exist as a single isomer.

Artificial amino acids with bulky side chains can be prepared according to known literature methods. For example, both L- and D- amino acid derivatives can be prepared starting directly from the commercially available L- or D-serine, respectively. Using this method, alkyl, phenyl and other bulky groups can be incorporated into serine to produce β -hydroxy- α -amino acids. Hence, artificial amino acids with bulky side chains can be incorporated into peptidic chelators, which would produce a single species and a single isomeric metal complex.

The advantage of having a bifunctional chelator that forms a single isomeric metal complex is that in the labeling of biologically important molecules, there is only a single radiolabeled species. Hence, there is no need to isolate and evaluate the biological activity and toxicity of multiple compounds. It is also easier to formulate a radiopharmaceutical kit that consistently produces a single radiolabeled compound than one that produces a series of radiolabeled compounds. In the labeling of a biologically important molecule with a chelator that results in multiple species, there is a necessity to formulate the kit such that the labeling consistently produces the same set of compounds in the same ratio. This is eliminated with the use of a chelator that produces a single metal complex. Quality control of a radiopharmaceutical is also simplified by the use of a chelator that results in a single species as it is much easier to develop a quality control protocol that identifies a single well characterized compound than one that has to identify the presence and quantity of multiple compounds.

An additional benefit of the incorporation of different side chain groups into the peptidic chelator backbone to cause a single isomer is that the lipophilicity of the resulting metal complexes is altered by the addition of the different groups. The log D of the ^{99m}Tc complex of dimethylglycine-L-t-butylglycine-L-cysteine-glycine is -1.3, compared to -2.3 for the ^{99m}Tc complex of dimethylglycine-L-serine-L-cysteine-glycine.

The terms defining the variables $R^1 - R^{10}$, $R^a - R^n$ and X as used hereinabove in formula (I) have the following meanings:

"alkyl" refers to a straight or branched C₁-C₈ chain and includes lower C₁-C₄ alkyl;

"alkoxy" refers to straight or branched C₁-C₈ alkoxy and includes lower C₁-C₄ alkoxy;

"thiol" refers to a sulfhydryl group that may be substituted with an alkyl group to form a thioether;

"sulfur protecting group" refers to a chemical group that is bonded to a sulfur atom and inhibits oxidation of sulfur and includes groups that are cleaved upon chelation of the metal. Suitable sulfur protecting groups include known alkyl, aryl, acyl, alkanoyl, aryloyl, mercaptoacyl and organothio groups.

"Linking group" refers to a chemical group that serves to couple the targeting molecule to the chelator while not adversely affecting either the targeting function of the peptide or the metal binding function of the chelator. Suitable linking groups include alkyl chains; alkyl chains optionally substituted with one or more substituents and in which one or more carbon atoms are optionally replaced with nitrogen, oxygen or sulfur atoms. Other suitable linking groups include those having the formula A¹-A²-A³ wherein A¹ and A³ are independently selected from N, O and S; and A² includes alkyl optionally substituted with one or more substituents and in which one or more carbon atoms are optionally replaced with nitrogen, oxygen or sulfur atoms; aryl optionally substituted with one or more substituents; and heteroaryl optionally substituted with one or more substituents. Still other suitable linking groups include amino acids and amino acid chains functionalized with one or more reactive groups for coupling to the glycopeptide and/or chelator. In one embodiment, the linking group is a peptide of 1 to 5 amino acids and includes, for example, chains of 1 or more synthetic amino acid residues such as β-Alanine residues. In another embodiment, the linking group is NH-alkyl-NH.

"Targeting molecule" refers to a molecule that can selectively deliver a chelated radionuclide or MRI contrasting agent to a desired location in a mammal. Preferred targeting molecules selectively target cellular receptors, transport systems, enzymes, glycoproteins and processes such as fluid pooling. Examples of targeting molecules suitable for coupling to the chelator include, but are not limited to, steroids, proteins,

peptides, antibodies, nucleotides and saccharides. Preferred targeting molecules include proteins and peptides, particularly those capable of binding with specificity to cell surface receptors characteristic of a particular pathology. For instance, disease states associated with over-expression of particular protein receptors can be imaged by labeling that protein or a receptor binding fragment thereof coupled to a chelator of invention. Most preferably targeting molecules are peptides capable of specifically binding to target sites and have three or more amino acid residues. The targeting moiety can be synthesised either on a solid support or in solution and is coupled to the next portion of the chelator-targeting moiety conjugates using known chemistry.

Chelator conjugates of the invention may be prepared by various methods depending upon the chelator chosen. The peptide portion of the conjugate if present is most conveniently prepared by techniques generally established in the art of peptide synthesis, such as the solid-phase approach. Solid-phase synthesis involves the stepwise addition of amino acid residues to a growing peptide chain that is linked to an insoluble support or matrix, such as polystyrene. The C-terminus residue of the peptide is first anchored to a commercially available support with its amino group protected with an Nt-butyloxycarbonyl protecting agent such as a group (tBoc) fluorenylmethoxycarbonyl (FMOC) group. The amino protecting group is removed with suitable deprotecting agents such as TFA in the case of tBOC or piperidine for FMOC and the next amino acid residue (in N-protected form) is added with a coupling agent such as dicyclocarbodiimide (DCC). Upon formation of a peptide bond, the reagents are washed from the support. After addition of the final residue, the peptide is cleaved from the support with a suitable reagent such as trifluoroacetic acid (TFA) or hydrogen fluoride (HF).

Conjugates may further incorporate a linking group component that serves to couple the peptide to the chelator while not adversely affecting either the targeting function of the peptide or the metal binding function of the chelator.

In accordance with one aspect of the invention, chelator conjugates incorporate a diagnostically useful metal capable of forming a complex. Suitable metals include radionuclides such as technetium and rhenium in their various forms such as ^{99m}TcO₃⁺, ^{99m}TcO₂⁺, ReO₃⁺ and ReO₂⁺. Incorporation of the metal within the conjugate can be

achieved by various methods common in the art of coordination chemistry. When the metal is technetium-99m, the following general procedure may be used to form a technetium complex. A peptide-chelator conjugate solution is formed initially by dissolving the conjugate in aqueous alcohol such as ethanol. The solution is then degassed to remove oxygen then thiol protecting groups are removed with a suitable reagent, for example with sodium hydroxide and then neutralized with an organic acid such as acetic acid (pH 6.0-6.5). In the labelling step, a stoichiometric excess of sodium pertechnetate, obtained from a molybdenum generator, is added to a solution of the conjugate with an amount of a reducing agent such as stannous chloride sufficient to reduce technetium and heated. The labelled conjugate may be separated from contaminants ^{99m}TcO₄ and colloidal ^{99m}TcO₂ chromatographically, for example with a C-18 Sep Pak cartridge.

In an alternative method, labelling can be accomplished by a transchelation reaction. The technetium source is a solution of technetium complexed with labile ligands facilitating ligand exchange with the selected chelator. Suitable ligands for transchelation include tartarate, citrate and heptagluconate. In this instance the preferred reducing reagent is sodium dithionite. It will be appreciated that the conjugate may be labelled using the techniques described above, or alternatively the chelator itself may be labelled and subsequently coupled to the peptide to form the conjugate; a process referred to as the "prelabelled ligand" method.

Another approach for labelling conjugates of the present invention involves techniques described in a co-pending United States application 08/152,680 filed 16 November 1993, incorporated herein by reference. Briefly, the chelator conjugates are immobilized on a solid-phase support through a linkage that is cleaved upon metal chelation. This is achieved when the chelator is coupled to a functional group of the support by one of the complexing atoms. Preferably, a complexing sulfur atom is coupled to the support which is functionalized with a sulfur protecting group such as maleimide.

A conjugate labelled with a radionuclide metal such as technetium-99m may be administered to a mammal by intravenous injection in a pharmaceutically acceptable solution such as isotonic saline. The amount of labelled conjugate appropriate for administration is dependent upon the distribution profile of the chosen conjugate in the

sense that a rapidly cleared conjugate may be administered in higher doses than one that clears less rapidly. Unit doses acceptable for imaging inflammation are in the range of about 5-40 mCi for a 70kg individual. In vivo distribution and localization is tracked by standard scintigraphic techniques at an appropriate time subsequent to administration; typically between 30 minutes and 180 minutes depending upon the rate of accumulation at the target site with respect to the rate of clearance at non-target tissue.

List of Abbreviations

Abbreviation	Description				
Acm	acetoamidomethyl				
Ar	argon				
Arg	arginine				
Boc	tert-butyloxycarbonyl				
Cys	cysteine				
DIEA	diisopropylethylamine				
Dimethylgly	N,N-dimethylglycine				
DMF	N,N-dimethylformamide				
ES-MS	Electron Spray Mass Spectrometry				
Fmoc	9-fluorenylmethyloxycarbonyl				
Gly	glycine				
HBTU	2-(1H-benzotriazol-1-yl)-1,1,3,3-tetramethyl-uronium hexafluorophosphate				
HOBT	1-hydroxybenzotriazole				
HPLC	high performance liquid chromatography				
Ile	isoleucine				
Leu	leucine				
Lys	lysine				
mL	millilitre(s)				
mmol	millimole(s)				
mol	mole(s)				
Mott	4-methoxytrityl				
NaOH	sodium hydroxide				
NMP	N-methylpyrrolidone				

Phe

phenylalanine

Pmc

2,2,5,7,8-pentamethylchroman-6-sulfonyl

 R_{t}

retention time

sasrin

2-methoxy-4-alkoxybenzyl alcohol (super acid sensitive

resin)

Ser

serine

t-Bu

tert-butyl

TFA

trifluoroacetic acid

Thr

threonine

Trt

trityl

Tyr

tyrosine

Yε-R

Y,

protection group R is attached to the peptide chain via the atom,

on the amino acid side chain (Y is N, O or S and R is

Acm, Boc,

Mott, t-Bu or Trt)

Examples

Materials. N-methylpyrrolidone, N,N-dimethylformamide, 100 mmol 2-(1H-benzotriazol-1-yl)-1,1,3,3-tetramethyl-uronium hexafluorophosphate/ 0.5M 1-hydroxybenzotriazole DMF, 2.0M diisopropylethylamine/ NMP, dichloromethane and trifluoroacetic acid were purchased from Applied Biosystems Inc. Triethylamine and *tert*-butyl methyl ether were purchased from Aldrich Chemical Inc. Fmoc amino acid derivatives and Fmoc-Gly sasrin resin was purchased from Bachem Bioscience Inc. All chemicals were used as received. [ReO₂(en)₂]Cl was prepared according to literature methods.^{57,58}

Instrumentation. NMR spectra were recorded on a Bruker AC-300 and on a Bruker DRX-500 NMR spectrometer and are reported as δ in ppm from external TMS. Mass spectra (electrospray) were obtained on a Sciex API#3 mass spectrometer in the positive ion detection mode. HPLC analyses and purifications were made on a Beckman System Nouveau Gold chromatographic system with a Waters 4 mm radial pak C-18 column. During analytical HPLC analysis, the mobile phase was changed from 100% 0.1% aqueous trifluoroacetic acid to 100% acetonitrile containing 0.1% trifluoroacetic acid over 20 minutes at a flow rate of 2 mL/min. All HPLC analyses were monitored with a UV detector set at 214 and 254 nm. Solid phase peptide syntheses were performed on an ABI Peptide Synthesizer model 433A using FastMoc chemistry and preloaded Fmoc amino acid sasrin resin. 59,60 Molecular modeling of the Re complexes was performed using Quanta Charm version 3.3.63 HPLC analyses of the 99mTc samples were made on a Beckman System Gold chromatographic system with a Vydac 4.6 mm radial pak C-18 column. The mobile phase was changed from 100% water containing 0.1% trifluoroacetic acid to 70% acetonitrile containing 0.1% trifluoroacetic acid over 25 minutes at a flow rate of 1 mL/min. The HPLC analyses of the 99mTc samples were monitored with a UV detector set at 215 nm and a radiometric gamma detector.

Example 1

Synthesis of Peptides. Peptides of various amino acid sequences were prepared via a solid phase peptide synthesis method on an automated peptide synthesizer using

FastMoc 1.0 mmole chemistry. ^{59,60} Preloaded Fmoc amino acid sasrin resin and Fmoc amino acid derivatives were used. Prior to the addition of each amino acid residue to the N-terminus of the peptide chain, the FMOC group was removed with 20% piperidine in NMP. Each Fmoc amino acid residue was activated with 0.50 M HBTU/ HOBt/ DMF, in the presence of 2.0M DIEA/ NMP. The C-terminus of the completed peptide was attached to the resin via the sasrin linker. The peptidyl resin was washed with dichloromethane and dried under vacuum for 20-24 hours. The peptide was cleaved off the resin by stirring the peptidyl resin in 95 % aqueous trifluoroacetic acid for 3-4 hours. The sasrin resin was filtered and the filtrate was added dropwise to *tert*-butyl methyl ether at 0 °C. The peptide precipitate out of the ether. The precipitate was collected by centrifugation and dissolved in minimal amount of water. The aqueous peptide solution was lyophilized to yield the product. The product was analyzed by mass spectrometry and by HPLC. The product was purified by HPLC. This method was used to prepare the following peptides

1)RP349: Dimethylgly-L-Ile-L-Cys(Sε-Acm)-Gly

2)RP332: Dimethylgly-L-lysine(N^ε-Biotin)-L- Cys(S^ε-Acm)

3)RP455: Dimethylgly-L-t-Butylgly-L-Cys(Sε-Acm)-Gly

4)RP505: Dimethylgly-D-t-Butylgly-L-Cys(Sε-Acm)-Gly

5)RP502: Dimethylgly-L-t-Butylgly-L-Cys(S^E-Acm)-Gly-Thr-Lys-Pro-Pro-Arg

6)RP573: Dimethylgly-L-t-Butylgly-L-Cys(S^e-Acm)-Gly-Arg-Ile-Lys-Pro-His

Example 2

Synthesis of Re Oxo Complex of Dimethylglycine-L-*t***-butylgly-L-Cys-Gly:** To remove the acm protecting group, dimethylgly-L-*t*-butylgly-L-Cys-(S^e-Acm)-Gly (84.0 mg, 0.187 mmoles) was dissolved in 2 mL of 30% acetic acid. Mercury(II) acetate (91.6 mg, 0.287 mmoles) was added to the solution and the solution was stirred under Ar at room temperature for 18 hours. H₂S gas was then bubbled through the solution for 5 minutes, causing black HgS to precipitate. The precipitate was removed by centrifugation, and the filtrate was frozen and lyophilized overnight. [ReO₂(en)₂]Cl (88.6 mg, 0.237 mmoles) was dissolved in 3 mL of distilled water and added to the lyophilized deprotected peptide. The solutions was a light green colour.

The pH of the solution was adjusted to 6 using 1 M NaOH. The solution was refluxed under Ar for 2 hours, during which time the solution changed from green to red. The solution was frozen and lyophilized overnight, yielding a red solid. Purification of the product was done by HPLC. Mass spectrum (electrospray): m/z = 577 ([MH]⁺), [C₁₅H₂₇N₄O₆Re₁S₁]. HPLC retention time: 9.52 min. ¹H NMR and ¹³C NMR (500 MHz, D₂O) spectral data are shown in Table 3 and 4. Log D (pH: 7.4): -1.3.

Example 3

Synthesis of Re Oxo Complex of Dimethylgly-D-t-butylgly-L-Cys-Gly: The procedure for the synthesis of the Re oxo complex of dimethylgly-D-t-butylgly-L-Cys-Gly was the same as the one described for the synthesis of the Re complex of Dimethylgly-L-t-butylgly-L-Cys-Gly. Mass spectrum (electrospray): m/z = 577 ([MH]⁺), [C₁₅H₂₆N₄O₆Re₁S₁]. HPLC retention time: 9.62 min. ¹H NMR (300 MHz, D₂O): 2.89 (s, methyl ¹H in the dimethylglycine residue), 3.65 (s, methyl ¹H in the dimethylglycine residue).

Example 4

Synthesis of Re Oxo Complex of Dimethylgly-L-t-Butylgly-L-Cys-Gly-Thr-Lys-Pro-Pro-Arg: The procedure for the synthesis of the Re oxo complex Dimethylgly-L-t-Butylgly-L-Cys-Gly-Thr-Lys-Pro-Pro-Arg was the same as the one described for the synthesis of the Re complex of dimethylgly-L-t-butylgly-L-Cys-Gly. Mass spectrum (electrospray): m/z = 1155 ([MH]⁺), [C₄₁H₇₁N₁₃O₁₂Re₁S₁]⁺). HPLC retention time: 8.82 min. ¹H NMR (500 MHz, D₂O): 2.63 (s, methyl ¹H in the dimethylglycine residue), 3.56 (s, methyl ¹H in the dimethylglycine residue).

Example 5

Synthesis of Re Oxo Complex of Dimethylgly-L-Ile-L-Cys-Gly: The procedure for the synthesis of the Re oxo complex Dimethylgly-L-ile-L-cys-gly was the same as the one described for the synthesis of the Re complex of dimethylgly-L-t-butylgly-L-cys-gly. Mass spectrum (electrospray): m/z = 577 ([MH]⁺), $[C_{41}H_{71}N_{13}O_{12}Re_1S_1]^+$), m/z = 598 ([MH]⁺, $[C_{41}H_{71}N_{13}O_{12}Re_1S_1]^+$). HPLC retention time: 9.50 min. ¹H NMR (300 MHz, D₂O): 2.60 (s, methyl ¹H in the dimethylglycine residue of isomer A), 2.76 (s,

methyl ¹H in the dimethylglycine residue of isomer B), 3.68 (s, methyl ¹H in the dimethylglycine residue of isomer A), 3.72 (s, methyl ¹H in the dimethylglycine residue of isomer B).

Example 6

Synthesis of the Re Oxo Complex of Dimethylgly-L-t-Butylgly-L-Cys-Gly-Arg-Ile-Lys-Pro-His: The procedure for the synthesis of the Re oxo complex of dimethylgly-L-t-Butylgly-L-Cys-Gly-Arg-Ile-Lys-Pro-His was the same as the one described for the synthesis of the Re complex of dimethylgly-L-t-butylgly-L-Cys-Gly. Mass spectrum (electrospray): m/z = 1207 ([MH]⁺), [C₄₃H₇₁N₁₅O₁₀Re₁S₁]. HPLC retention time: 8.78 min. ¹H NMR (300 MHz, D₂O): 2.71 (s, methyl ¹H in the dimethylglycine residue).

Example 7

Synthesis of the ^{99m}Tc **complex**. The peptide (0.2-0.5 μmoles) was dissolved in 200 μL of saline. Na[^{99m}TcO₄] (10 mCi) was added to the solution, followed by tin(II) chloride (7.5 x 10³ μg, 39 μmoles), sodium gluconate (1.3 x 10³ μg, 5.8 μmoles), and 20 μL of 0.1 M NaOH. The solution was left at room temperature for 1 hour or heated at 100 °C for 15 minutes. In the synthesis of the ^{99m}Tc complex, the acetoamidomethyl protection group was displaced from the cysteine residue in RP414. The ^{99m}Tc complex was analyzed by HPLC. The ^{99m}Tc complexes of RP455, RP505 and RP502 was co-injected with the corresponding Re complexes. The HPLC retention times of the ^{99m}Tc peptidic complexes are as follows:

- 1)^{99m}Tc complex of RP349 (Dimethylgly-L-Ile-L-Cys-Gly): HPLC retention time: ^{99m}Tc(RP349) R₄ = 19.41, 21.53 min (radiometric gamma detector).
- 2)^{99m}Tc complex of RP332 (Dimethylgly-L-lysine(N^ε-Biotin)-L- Cys): HPLC retention time: ^{99m}Tc(RP332) R_s = 11.54, 11.97 min (radiometric gamma detector).
- 3)^{99m}Tc complex of RP455 (Dimethylgly-L-t-Butylgly-L-Cys-Gly): HPLC retention time: ReO(RP455) R_t = 21.18 min (UV detector set at 215 nm); ^{99m}Tc(RP445) R_t = 21.49 min (radiometric gamma detector).

4) ^{99m}Tc complex of RP505 (Dimethylgly-D-*t*-Butylgly-L-Cys-Gly): HPLC retention time: ReO(RP505) R_t = 18.16 min (UV detector set at 215 nm); ^{99m}Tc(RP505) R_t = 18.89 min (radiometric gamma detector).

- 5)^{99m}Tc complex of RP502 (Dimethylgly-L-t-Butylgly-L-Cys(S $^{\epsilon}$ -Acm)-Gly-Thr-Lys-Pro-Pro-Arg): HPLC retention time: ReO(RP502) R_t = 19.76 min (UV detector set at 215 nm); ^{99m}Tc(RP502) R_t = 20.10 min (radiometric gamma detector).
- 6) ^{99m}Tc complex of RP573 (Dimethylgly-L-t-Butylgly-L-Cys(S^ε-Acm)-Gly-Arg-Ile-Lys-Pro-His): HPLC retention time: (ReORP573) R_t = 16.43 min (UV detector set at 215 nm); ^{99m}Tc(RP573) Rt = 20.75 min (radiometric gamma detector).

Example 8

Synthesis of Dimethylgly-L-Beta-hydroxyvaline-L-Cys-Gly. The beta-

hydroxyvaline is synthesized according to the method of Shao, H., and Goodman, M., *J. Org. Chem.* **1996**, *61*, 2582-2583 or Beloken, Yu. N.; Bulychev, A. G.; Vitt, S. V.; Struchkov, Yu. T.; Batsanov, A. S.; Timofeeva, T. V.; Tsyryapkin, V. A.; Ryzhov, M. G.; Lysova, L. A.; *Et al. J. Am. Soc. Chem.*, **1985**, *107(14)*, 4252-9. The FMOC group is added to the amino terminus according to the method of Carpino, L. A., Han, G. Y. *J. Org. Chem.* **1972**, *37*, 3404. The FMOC-beta-hydroxyvaline is purified by column chromatography. The peptide dimethylgly-L-beta-hydroxyvaline-L-cys-gly is synthesized on the peptide synthesizer in the same manner as set out in Example 1. The Re and Tc-99m complexes are synthesized by the same method as the Re and Tc-99m complexes of dimethylgly-L-t-butylgly-L-cys-gly as shown in examples 2 and 7 respectively.

These Re and Tc-99m chelates form the syn isomer predominantly but are more hydrophilic than the Re and Tc-99m complexes already mentioned. This is an advantage when the chelates are attached to hydrophilic targeting molecules.

Although the invention has been described with preferred embodiments, it is to be understood that modifications may be resorted to as will be apparent to those skilled

in the art. Such modifications and variations are to be considered within the purview and scope of the present invention.

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We Claim:

1. A chirally pure compound of the formula I:

$$R^{1}$$
 R^{3}
 R^{4}
 R^{5}
 R^{10}
 R^{10}
 R^{10}
 R^{10}
 R^{10}
 R^{10}
 R^{10}
 R^{10}

wherein

R¹ is a linear or branched, saturated or unsaturated C_{1.4}alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents selected from halogen, hydroxyl, amino, carboxyl, C_{1.4}alkyl, aryl and C(O)R¹⁰;

R² is H or a substituent defined by R¹;

- R¹ and R² may together form a 5- to 8-membered saturated or unsaturated heterocyclic ring optionally substituted by one or more substituents selected from halogen, hydroxyl, amino, carboxyl, oxo, C₁₋₄alkyl, aryl and C(O)Z;
- R³, R⁴ and R⁵ are selected independently from H; carboxyl; C₁₋₄alkyl; C₁₋₄alkyl substituted with a substituent selected from hydroxyl, amino, sulfhydryl, halogen, carboxyl, C₁₋₄alkoxycarbonyl and aminocarbonyl; an alpha carbon side chain of a D- or L-amino acid other than proline; and C(O)R¹⁰;

R⁶ is selected from the group consisting of:

- i) an optionally substituted 3- to 6-membered heterocylic or carbocylic ring,;
- ii) a compound having the following formula:

wherein R^{11} , R^{12} and R^{13} are independently selected from H, linear or branched, saturated or unsaturated C_{1-6} alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents, alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R^{11} , R^{12} and R^{13} is not H;

iii) a compound of the following formula:

wherein R^{14} and R^{15} are independently selected from H, linear or branched, saturated or unsaturated $C_{1.6}$ alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R^{14} and R^{15} is not H; and

iv) a compound of the following formula:

wherein X is selected from O or S and R¹⁶ is selected from linear or branched, saturated or unsaturated C₁₋₆alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents (; alkoxycarbonyl, aminocarbonyl, alkoxy, and an optionally substituted 3- to 6-membered heterocylic or carbocylic ring;

R⁷ and R⁸ are selected independently from H; carboxyl; amino; C₁₋₄alkyl; C₁₋₄alkyl substituted by a substituent selected from hydroxyl, carboxyl and amino; and C(O)R¹⁰;

R9 is selected from H and a sulfur protecting group; and

R¹⁰ is selected from hydroxyl, alkoxy, an amino acid residue, a linking group and a targeting molecule.

2. A chirally pure compound of the formula II:

wherein

R^a is selected from H and a sulfur protecting group;

 R^b , R^c R^d , R^f and R^g are selected independently from H; carboxyl; C_{1-4} alkyl; C_{1-4} alkyl substituted with a substituent selected from hydroxyl, amino, sulfhydryl, halogen, carboxyl, C_{1-4} alkoxycarbonyl and aminocarbonyl; an alpha carbon side chain of a D- or L-amino acid other than proline; and $C(O)R^h$;

R^e is an optionally subsituted 3- to 6-membered heterocylic or carbocylic ring; or R^e is

wherein Rⁱ, R^j and R^k are independently selected from H, linear or branched, saturated or unsaturated C₁₋₆alkyl chain that is optionally interrupted by one or

two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of Rⁱ, R^j and R^k is not H;

or Re is

wherein R¹ and R^m are independently selected from H, linear or branched, saturated or unsaturated C₁₋₆alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; with the proviso that a least one of R¹ and R^m is not H;

or Re is

wherein X is selected from O or S and Rⁿ is selected from linear or branched, saturated or unsaturated C_{1.6}alkyl chain that is optionally interrupted by one or two heteroatoms selected from N, O and S; and is optionally substituted by one or more substituents; alkoxycarbonyl, aminocarbonyl, alkoxy, and an optionally substituted 3- to 6-membered heterocylic or carbocylic ring; and

R^h is selected from hydroxyl, alkoxy, an amino acid residue, a linking group and a targeting molecule.

3. A compound selected from:

Dimethylgly-L-t-Butylgly-L-Cys-Gly;

Dimethylgly-D-t-Butylgly-L-Cys-Gly;

Dimethylgly-L-t-Butylgly-L-Cys; and

Dimethylgly-L-t-Butylgly-L-Cys(S $^{\epsilon}$ -Acm)-Gly-Thr-Lys-Pro-Pro-Arg.

4. A compound accrding to any of claims 1 to 3 in a from complexed with a metal or metal radionuclide or an oxide or nitride thereof.

- 5. A pharmaceutical composition, comprising a pharmaceutically acceptable carrier and a compound as defined in claim 4 in an amount effective to image a site of diagnostic interest.
- 6. A method of radioimaging a site of diagnostic interest, comprising the step of administering systemically to a patient a pharmaceutical composition as defined in claim 5, allowing the pharmaceutical to localize within the site of diagnostic interest, and then taking an image of the patient so treated.

INTERNATIONAL SEARCH REPORT

Internat I Application No PCT/CA 98/01201

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Name and	mailing address of the ISA European Patent Office, P.B. 5818 Patentlaan 2	Authorized officer			
	European Patent Office, P.B. 3010 Patermaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni,	Kronester-Frei, /	1		
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